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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Title:
Perfect score:
                                                                                                                                                                                                                                                                 Result
                                                                                                                                                                                                                                                                                                                                                                   Database
                                                                                                                                                                                                                                                                                                                                                                                                                                          Total number of hits satisfying chosen parameters:
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  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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294
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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ALIGNMENTS

apoptosis inhibitor hiap-1 - human C;Species: Homo sapiens (man) C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 21-Jul-2000 C;Accession: 568449

R; Liston, P.; Roy, N.; Tamai, K.; Lefebvre, C.; Baird, S.; Cherton-Horvat, G., Fur.P., Nature 379, 349-353, 1996
A;Title: Suppression of apoptosis in mammalian cells by NAIP and a related family c., A;Reference number; A58182; MUID:96149249

A;Status: nucleic acid sequence not shown A;Molecule type: mRNA A;Residues: 1-604 <LIS>

A: Accession: S68449

RESULT S68449

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C;Accession: UC5964
R;Stchilk, C; de Martin, R; Binder, B.R; Lipp, J. Biochem. Blophys. Res. Commun. 243, 827-832, 1998
A;Title: Cytokine induced expression of porcine inhil A;Reference number: UC5964; MUID:98162622
A;Accession: UC5964
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-358 <STE>
                                                                                                                                                                                                                                                                    apoptosis inhibitor - pig
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 04-Sep-1998 #sequence_revision 04-Sep-1998 #text_change 02-Sep-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A:Description: apoptotic suppressor
C:Superfamily: RING finger homology
C:Reywords: apoptosis; zinc finger
C:Reywords: RING finger homology <RNG>
A;Cross-references: GB:U79142; NID:g2957174; C;Superfamily: RING finger homology F;307-351/Domain: RING finger homology <RRN>
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Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   273 PEQLASAGFYYYGNSDDVKCFCCDGGLRCWESGDDPWVQHAKWFPRCE 320
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Pred. No. 4.3e-27;
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                                               PIDN:AAC39171.1; PID:g2957175
                                                                                                                                                                                       inhibitor of apoptosis protein (iap)
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Query Match Best Local Similarity

96.3%;

Score 283; DB 2; pred. No. 5.3e-26;

Length 358;

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apoptosis-inhibiting protein - Orgyia pseudotsugata multicapsid nuclear polyhedrosis C:Species: Orgyia pseudotsugata multicapsid nuclear polyhedrosis virus, OpMNPV C:Date: 15-Oct-1994 #sequence_revision 15-Oct-1994 #text_change 15-Sep-2000 C:Accession: A53989 #Sequence_revision 15-Oct-1994 #text_change 15-Sep-2000 R;Birnbaum, M.J.; Clem. B 7 · uilland R;Birnbaum, M.J.; Cl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inhibitor of apoptosis protein 3 - Orgyla pseudotsugata nuclear polyhedrosis C;Specles: Orgyla pseudotsugata nuclear polyhedrosis virus, OpNNPV C;Date: 16-Vul-1999 #sequence_revision 16-Vul-1999 #text_change 15-Sep-2000 C;Accession: T10304 R;Ahrens - C.A.; Russell, R.R.; Funk, C.J.; Evans, J.; Harwood, S.; Rohrmann, Virology 229, 381-399, 1997 R;Ahrens - C.A.; Russell, B. R. Speudotsugata multinucleocapsid nuclear F A;Reference number: Z17011; MUID:97Z71300 A;Accession: T10304 A;Accession: T10304
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C:Accession: P: Roy, N.: Tamai, K.; Lefebvre, C.;
Nature 379, 349-353, 1996
A:Title: Suppression of apoptosis in mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: DNA
A;Residues: 1-268 <AHR>
A;Residues: EBMB: U75930; NID:g2934903;
A;Cross-references: EBMB:U75930; NID:g2934903;
C;Superfamily: viral apoptosis inhibitor IAP; R
F;217-261/Domain: RING finger homology <RRN>
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C;Species: Homo sapiens (man)
C;Species: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 21-Jul-2000
C;Date: 17-Jul-1998
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A; Residues: 1-618 <LIS>
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Pred. No. 2.5e-16;
6; Mismatches 12
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Pred. No. 3e-24;
2; Mismatches
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    Query Match
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A. Nolecule type: DNA
A. Molecule type: DNA
A. Molecule type: DNA
A. Residues: 1-275 <CRO2
A. Note: sequence extracted from NCBI backbone (NCBIN:127014, NCBIP:127015)
A. Note: sequence extracted from NCBI backbone (NCBIN:127014, NCBIP:127015)
A. Note: sequence extracted from NCBI backbone (NCBIN:127014, NCBIP:127015)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Gene: kiap
A;Map position: 20q13
C;Keywords: apoptosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         J. Virol. 68, 2521-2528, 1994
A;Tille: An apoptosis inhibiting gene from a nuclear polyhedrosis virus
A;Reference number: A53989; MUID:94187094
A;Accession: A53989
A;Status: preliminary
                                                                                                                                                                                        R;Crook, N.E.; Clem, R.J.; Miller, L.K.
J. Virol. 67, 2168-2174, 1993
A;Title: An apoptosis-inhibiting baculovirus
A;Reference number: A45679; MUID:93188168
                                                                                                                                                                                                                                                                                              inhibitor-of-apoptosis polypeptide (IAP) - Cydia pomonella
C:species: Cydia pomonella granniosis virus CPGV
C:Date: 21-Sep-1993 #sequence_revision 25-Apr-1997 #text_cl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Reference number: JC7568; A; Contents: Fetal kidney A; Accession: JC7568
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R;Lin, J.H.; Deng, G.; Huang, Q.; Morser, J.
Biochem. Bloophys. Res. Commun. 279, 820-831, 2
A;Title: KIAP, a novel member of the inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: GB:L22564; NID:9456111; PIDN:AAB02610.J; PID:9456114 A;Note: authors translated the codon TGG for residue 28 as Tyr, GAC for ;C;Superfamilly: viral apoptosis! inhibitor IAP; RING finger homology F;217-261/Domain: RING finger homology <RRN>
                                                                                                                                            A; Status: preliminary
                                                                                                                                                                   A; Reference number: A45679; A; Accession: A45679
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A; Residues: 1-268 <BIR>
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Pred. No. 1.Be-15;
9; Mismatches 10
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A:Cross references: EMBL:U32974; NID:g1016687; PIDN:AAC50518.1; PID:g1016688 R;Liston, P; Roy, N: Tamal, K.; Lefebvre, C.; Baird, S.; Cherton-Horvat, G.; Nature 379, 349-353, 1996
A:Title: Suppression of apoptosis in mammalian cells by NAIP and a related fami A:Reference number: A58182; MUID:96149249
A:Accession: 868451
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                                                                                                                                                           A;Cross-references:
C;Genetics:
A;Gene: Naip2
                                                                                                                                                                                                                                                                                                                                                                                                               R;Yaraghi, 2.; Diez, E.; Gros, P.; MacKenzie, Mamm. Genome 10, 761-763, 1999
A;Title: CDNA cioning and the 5'genomic organ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neuronal apoptosis inhibitory protein 2 - mouse C;Species: Mus musculus (house mouse) C;Dete: 11-Jan-2000 #sequence_revision 11-Jan-2000 C;Accession: T42628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Description: apoptotic suppressor
C; Superfamily: apoptosis inhibitor IAP homolog;
C; Reywords: apoptosis; zinc finger
C; Reywords: apoptosis; zinc finger
F; 446-490/Domain: RING finger homology RRN>
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A;Residues: 1-161,'C',163-422,'Q',424-497 <LIS>
A;Cross-references: EMBL:U45880; NID:g1184319; PIDN:AAC50373.1; PID:g1184320
C;GeneriLs:
A;Gene: ilp
C;Function:
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A; Residues: 1-1447 < YAR>
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A; Accession: T42628
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A; Residues: 1-497 < DUC>
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5; Mismatches
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Pred. No. 2.2e-13;
                               Score 177; DB 2
Pred. No. 6e-13;
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                                                                                                                                                                                                                 R;Baird, S.D.

Submitted to the EMBL Data Library, January 1996

A;RcEarence number: $78528

A;RcEarence number: $78528

A;RcEarence rumber: $78528

C;Genetics: The content of the rumber rumb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   apoptosis inhibitor diap - fruit fly (Drosophila melanogaster) c;Species: Drosophila melanogaster c;Decies: Drosophila melanogaster c;Dete: 17-7ul-1998 #sequence_revision 17-Jul-1998 #text_chang c;Accession: 568452; $78528 R:Liston, P.; Royy, N.; Tamai, K.; Lefebvre, C.; Baird, S.; Che Nature 379, 349-353, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Title: The gene for neuronal apoptosis inhibitory protein is partially deleted A;Reference number: A55478; MUID:95112344
A;Recession: A55478
A;Recession: A55478
A;Residues: 1-1232 <ROY>
                           C;Function:
A;Description: apoptotic suppressor
C;Superfamily: apoptosis inhibitor IAP homolog;
C;Keywords: apoptosis; zinc finger
C;Keywords: apoptosis; zinc finger
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F;479-495/Domain: transmembrane #status predicted <MMM2>
F;479-496/Domain: transmembrane predicted <MMM2>
F;476/Binding site: ATP (Lys) #status predicted (Asn) (covalent) #status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: GB:U19251
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Roy, N.; Mahadevan, M.S.; Mod, T.O.; de Jong, P.J.; Surh, Cell 80, 167-178, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N; Alternate names:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           망
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A:PitLe: Suppression of apoptosis in mammalian cells by A;Reference number: A58182; MUID:96149249 A;Accession: $68492
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A;Map position: 5q12.2-5q13
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C; Date: 05-Jan-1996 #sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: mRNA
A; Residues: 1-496 <LIS>
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                                                                                                                                                                                              A;Cross-references: FlyBase:FBgn0015247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: EMBL:U45881;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Status: nucleic acid sequence not shown
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Best Local
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nes 28; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          K.; Lefebvre, C.; Baird, S.; Cherton-Horvat, G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         #status predicted <TMM1>
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Pred. No. 1.5e
5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NID:g1184313; PIDN:AAC46988.1; PID:g1184314
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleotide binding;
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.5e-12;
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T31067
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RESULT 14
T28409
T28409
ORF MSV248 probable inhibitor of apoptosis protein (IAP) - Melanoplus sanguinipes C;Species: Melanoplus sanguinipes entomopoxvirus
C;Species: Melanoplus sanguinipes entomopoxvirus
C;Date: 21-Jan-2000 #sequence_revision 21-Jan-2000 #text_change 21-Jul-2000
C;Accession: T28409
                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: EMBL:Y17267; NID:g3319989; PIDN:CAA76720.1; PID:g3319990 A;Note: localized to the Golgi compartment and the vesicular system C;Keywords: membrane-associated protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A:Pitle: A glant ubiquitin-conjugating erzyme related to IAP apoptosis inhibitors A:Reference number: 220963; MUID:98292517
A:Accession: 721067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BIR repeat containing ubiquitin-conjugating enzyme BRUCE - mouse C:Species: Mus musculus (house mouse) C:C:Date: 22-Oct-1999 #sequence_revision 22-Oct-1999 #text_change 21-Jul-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Accession: 56945
R;Duckett, C.S.; Nava, V.E.; Gedrich, R.W.; Clem, R.J.; van Dongen, J.L.; Gilfillan, M. EMBO J. 15, 2685-2694, 1996
A;Title: A conserved family of cellular genes related to the baculovirus iap gene and e A;Paterence number: 569544; MUID:96256286
A;Accession: 569545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                apoptosis inhibitor IAP homolog - fruit fly (Drosophila melanogaster) C:Species: Drosophila melanogaster C:Date: 06-Dec-1996 #sequence_revision 13-Mar-1997 #text_change 21-Jan-2000
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A;Molecule type: mRNA
A;Residues; 1-4845 <HAU>
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A; Residues: 1-497 < DUC>
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Best Local S
Matches 21
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                                                                                                                                                                                                            PEQLASAGFYY--VGNSDD-VKCFCCDGGLRCWESGDDPWVQHAKWFPRC 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LAQAGLYYOKIGDOVRCFHCNIGLRSWOKEDEPWFEHAKWSPKCQ 277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LAQAGLYYQKIGDQVRCFHCNIGLRSWQKEDEPWFEHAKWSPKCQ 275
                                                                                                                                                                                       PDPMAQAGFYHQPASSGDDRAMCFTCSVCLVCWEPTDEPWSEHERHSPNC 330
                                                                                                                                                                                                                                                                                                Similarity
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Pred. No. 1.5e-10;
7; Mismatches 14
                                                                                                                                                                                                                                                                                                   Score 116.5; DB 2
Pred. No. 2.5e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 153; DB 2;
Pred. No. 1.5e-10;
7; Mismatches 14
                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                        DB 2;
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4 LASAGFYYVGNSDDVKCFCCDGGLRCWESGDDPWVQHAKWFPRC 47

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38.8%;

Score 114; DB Pred. No. 3.6e 8; Mismatches

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IAEAGLEYTGRGDETVCFFCDCCVRDWHTNEDTWQRHAAENPQC

196

January

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15:49:11

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R;Afonso, C.L.; Tulman, E.R.; Lu, Z.; Oma, E.; Kutish, G.F.; Rock, D.L.
J. Virol. 73, 533-552, 1999
A;Title: The genome of Melanoplus sanguinipes entomopoxvirus.
A;Reference number: Z20484; MUID:99102612
A;Accession: T28409
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-150 (ARO)
A;Cross-references: EMBL:AF063866; NID:g4049647; PIDN:AAC97724.1; PID:g4.Note: MSV248
                                                                                                                                                                                               A;Residues: Î'286 <BRA>
A;Cross-references: GB:52569
A;Cross-references: GB:52569
R:Ayres, M.D.; HOward, S.C.; Kuzio, J.; Lopez-Ferber, M.; Possee, R.D.
Virology 202, 586-605, 1994
Virology 202, 586-605, 1994
A;Title: The complete DNA sequence of Autographa californica nuclear polyhedrosis vir
A;Reference number: A72850; MUID:94303173
A;Accession: C72853
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         orfil protein - Autographa californica nuclear polyhedrosis virus
C;Species: Autographa californica nuclear polyhedrosis virus, AcMNPV
C;Date: 30-Sep.1993 #sequencc_revision 30-Sep-1993 #text_change 17-Nov-2000
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                                                                                  A; Molecule type: DNA
A; Residues: 1-199, 'L', 201-286 <
A; Cross-references: GB:L22858;
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A;Gene: Ac-IAP1
C;Superfamily: viral apoptosis inhibitor IAP; RING finger homology
                                                             C; Genetics:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Virology 191, 1003-1008, 1992
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R; Braunagel, S.C.; Daniel,
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                              Ac-IAPI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .; Daniel, K.D.; Reilly, L.M.; Guarino, L.A.; Hong, T.; Summers,
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